AMENDMENTS

- (currently amended) A compound 12 to 50 nucleobases in length targeted to a region comprising nucleotide 901 to 950 of the nucleic acid molecule encoding diacylglycerol acyltransferase 2 in SEQ ID NO: 4, wherein said compound is at least 80% complementary to said nucleic acid molecule encoding diacylglycerol acyltransferase 2, and wherein said compound comprises at least an 8 nucleobase portion of SEO ID NO: 35, 36, 37 or 38.
- 2. (canceled)
- (previously presented) The compound of claim 1, wherein said compound is 15 to 30 nucleobases in length.
- 4. (original) The compound of claim 1 comprising an oligonucleotide.
- 5. (previously presented) The compound of claim 4 comprising an antisense oligonucleotide.
- 6. (original) The compound of claim 4 comprising a DNA oligonucleotide.
- 7. (original) The compound of claim 4 comprising a RNA oligonucleotide.
- 8. (original) The compound of claim 4 comprising a chimeric oligonucleotide.
- (original) The compound of claim 4 wherein at least a portion of said compound hybridizes with RNA to form an oligonucleotide-RNA duplex.
- 10. (canceled).
- 11. (original) The compound of claim 1 having at least 90% complementarity with said nucleic acid molecule encoding diacylglycerol acyltransferase 2.
- 12. (previously presented) The compound of claim 1 having at least 95% complementarity with said nucleic acid molecule encoding diacylglycerol acyltransferase 2.
- 13. (previously presented) The compound of claim 1 having 100% complementarity with said nucleic acid molecule encoding diacylglycerol acyltransferase 2.

- 14. (previously presented) The compound of claim 1 having at least one modified internucleoside linkage, sugar moiety, or nucleobase.
- 15. (original) The compound of claim 1 having at least one 2'-O-methoxyethyl sugar moiety.
- 16. (original) The compound of claim 1 having at least one phosphorothioate internucleoside linkage.
- 17. (original) The compound of claim 1 having at least one 5-methylcytosine.
- 18.-21. (canceled)
- 22. (original) A kit or assay device comprising the compound of claim 1.
- 23.-57. (canceled)
- 58. (previously presented) The compound of claim 1, wherein said compound is 20 nucleobases in length.
- 59. (previously presented) The compound of claim 13 having at least one modified internucleoside linkage, sugar moiety, or nucleobase.
- 60. (previously presented) A compound 20 nucleobases in length targeted to a nucleic acid molecule encoding diacylglycerol acyltransferase 2 (SEQ ID NO: 4), wherein said compound has the nucleobase sequence of SEQ ID NO: 35.

- 61. (previously presented) The compound of claim 60, wherein said compound is an antisense oligonucleotide comprising:
 - a gap segment consisting of linked deoxynucleosides;
 - a 5' wing segment consisting of linked nucleosides;
 - a 3' wing segment consisting of linked nucleosides;

wherein the gap segment is positioned between the 5' wing segment and the 3' wing segment and wherein each nucleoside of each wing segment comprises a modified sugar.

- 62. (previously presented) The compound of claim 61, wherein the antisense oligonucleotide comprises:
 - a gap segment consisting of ten linked deoxynucleosides;
 - a 5' wing segment consisting of five linked nucleosides;
 - a 3' wing segment consisting of five linked nucleosides;

wherein the gap segment is positioned between the 5' wing segment and the 3' wing segment, wherein each nucleoside of each wing segment comprises a 2'-O-methoxyethyl sugar; and wherein each internucleoside linkage is a phosphorothioate linkage.

- 63. (previously presented) The compound of claim 62, wherein each cytosine is a 5-methylcytosine.
- 64. (previously presented) The compound of claim 62, wherein the deoxynucleosides are 2'-deoxynucleosides.
- 65. (previously presented) The compound of claim 63, wherein the deoxynucleosides are 2'-deoxynucleosides.
- 66. (new) The compound of claim 1, wherein said compound comprises at least an 8 nucleobase portion of SEQ ID NO: 36.
- 67. (new) The compound of claim 1, wherein said compound comprises at least an 8 nucleobase portion of SEO ID NO: 37.
- 68. (new) The compound of claim 1, wherein said compound comprises at least an 8 nucleobase portion of SEQ ID NO: 38.

- 69. (new) A compound 12 to 50 nucleobases in length targeted to a region comprising nucleotide 901 to 950 of the nucleic acid molecule encoding diacylglycerol acyltransferase 2 in SEQ ID NO: 4, wherein said compound is 100% complementary to said nucleic acid molecule encoding diacylglycerol acyltransferase 2, and wherein said compound comprises the nucleotide sequence of SEO ID NO: 35.
- 70. (new) The compound of claim 69, wherein said compound is 15 to 30 nucleobases in length.
- 71. (new) The compound of claim 69 comprising an oligonucleotide.
- 72. (new) The compound of claim 71 comprising an antisense oligonucleotide.
- 73. (new) The compound of claim 71 comprising a DNA oligonucleotide.
- 74. (new) The compound of claim 71 comprising a RNA oligonucleotide.
- 75. (new) The compound of claim 71 comprising a chimeric oligonucleotide.
- 76. (new) The compound of claim 71 wherein at least a portion of said compound hybridizes with RNA to form an oligonucleotide-RNA duplex.
- 77. (new) The compound of claim 69 having at least one modified internucleoside linkage, sugar moiety, or nucleobase.
- 78. (new) The compound of claim 69 having at least one 2'-O-methox vethyl sugar mojety.
- 79. (new) The compound of claim 69 having at least one phosphorothioate internucleoside linkage.
- 80. (new) The compound of claim 69 having at least one 5-methylcytosine.
- 81. (new) A kit or assay device comprising the compound of claim 69.
- 82. (new) The compound of claim 69, wherein said compound is 20 nucleobases in length.